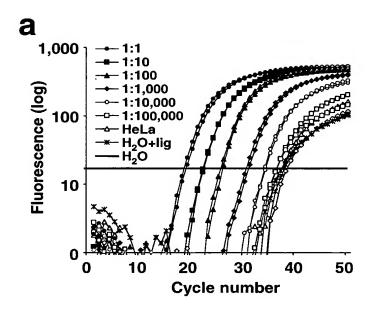
WT Wild-type DNA No amplification No ligation ↓Bottom strand synthesis VG-PCR probe cleavage Ligated DNA Mutant DNA Figure 1 **↓**Ligation **Mutant DNA** Ø Step 1. Ligation Step 2. Q-PCR

Figure 2



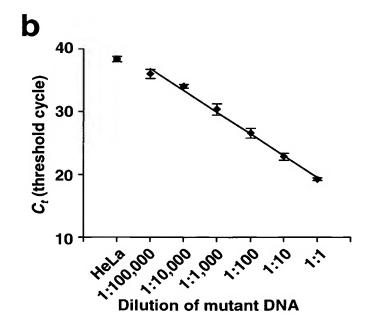
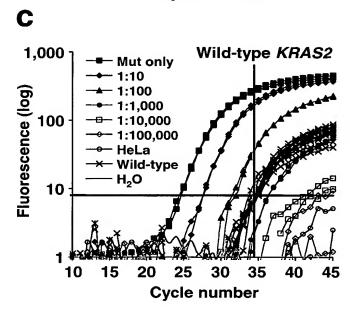


Figure 2 (cont.)

Cycle number



Dilution of mutant DNA

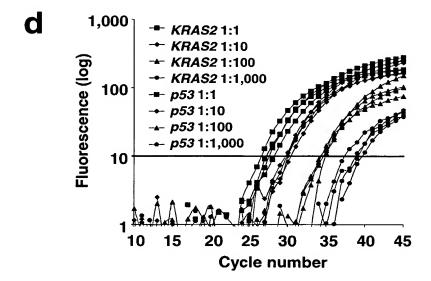


Figure 3

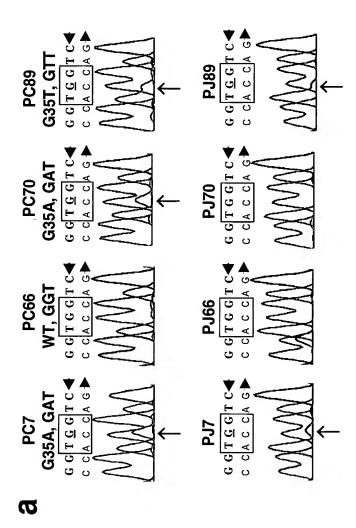
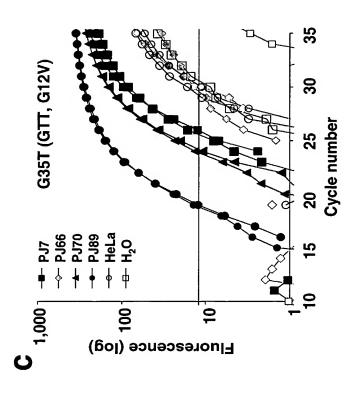


Figure 3 (cont.)



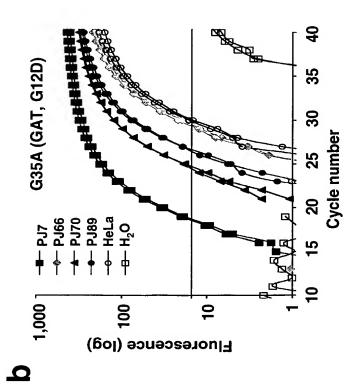
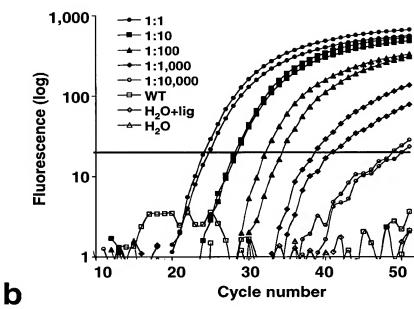


Figure 4





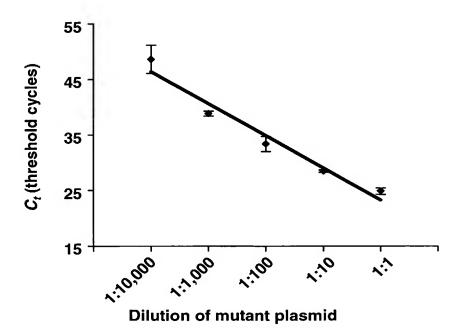
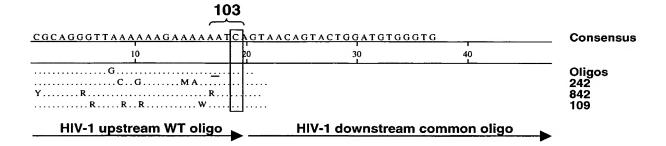
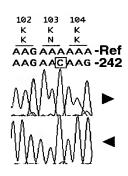


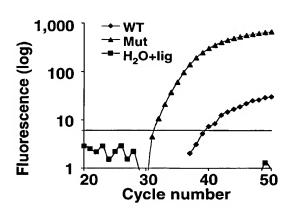
Figure 5

a



b





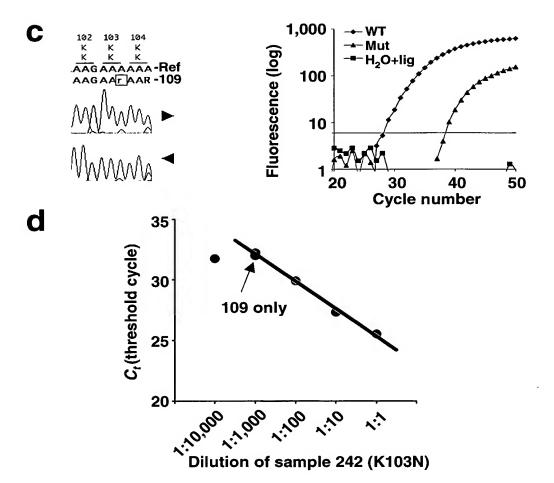


Figure 6

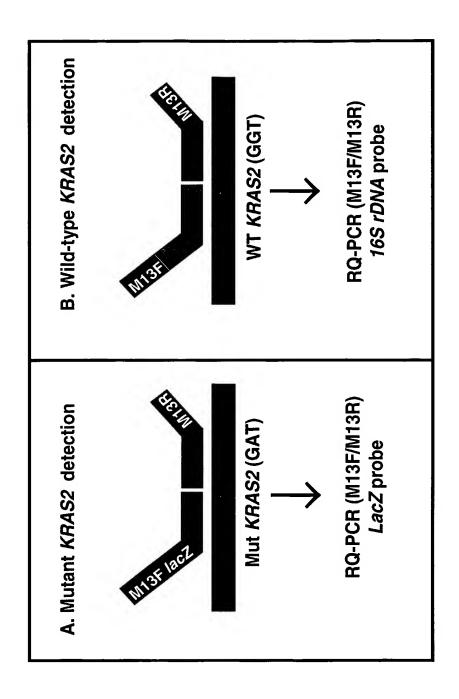


Figure 7A

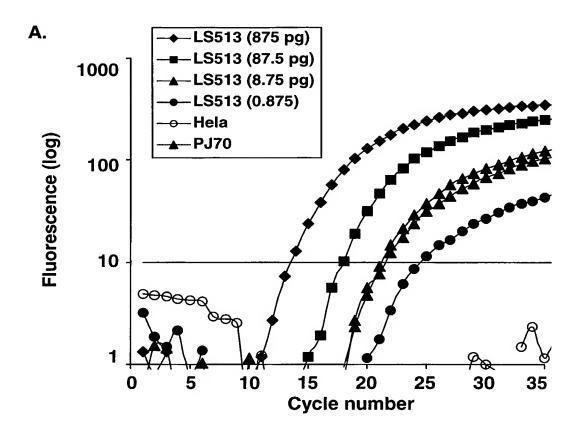
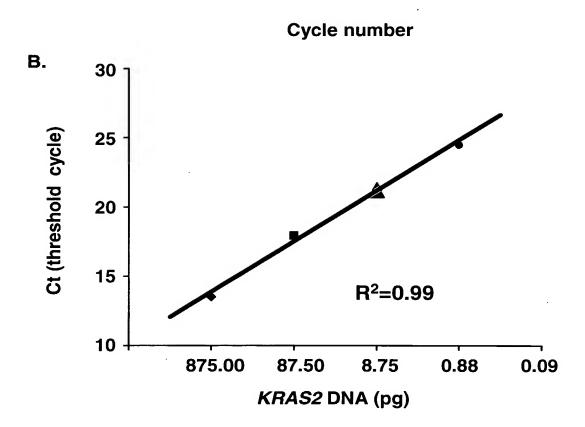
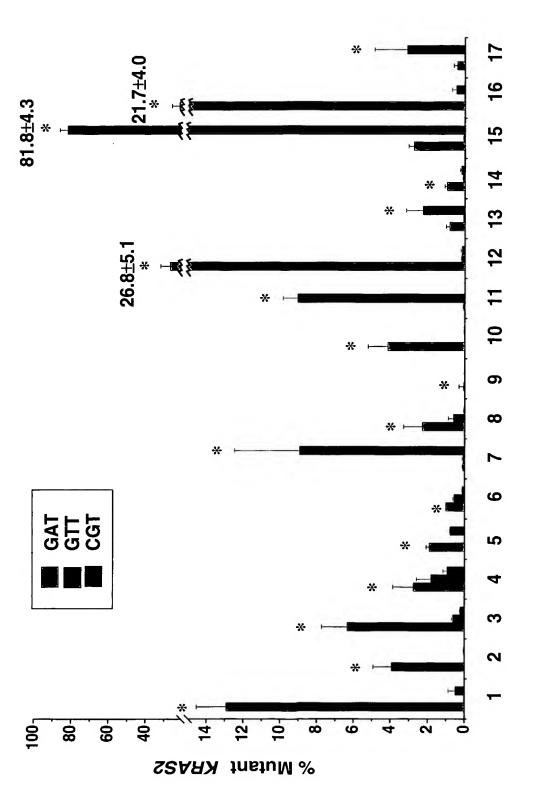


Figure 7B





WO 2005/092038 PCT/US2005/009391

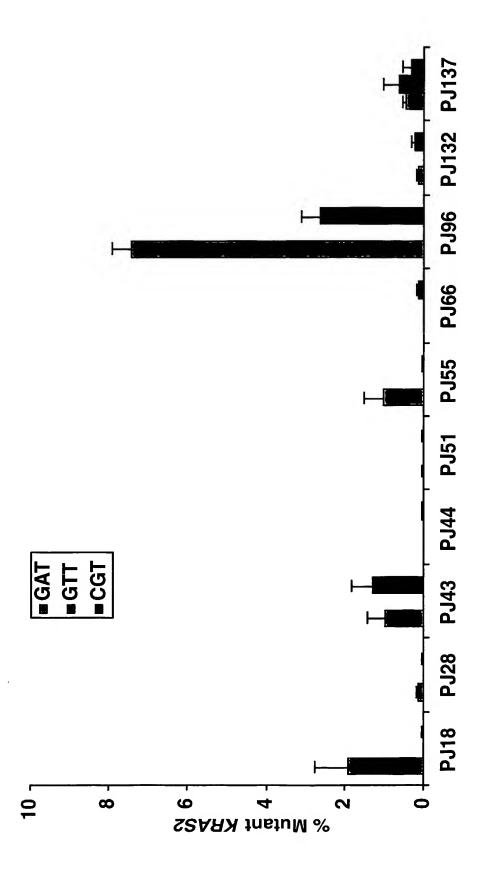


Figure 9

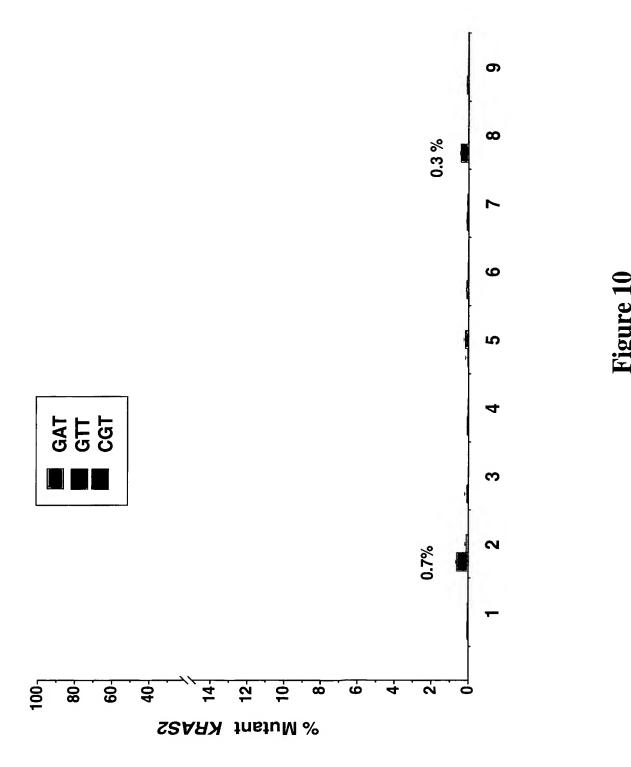


Figure 11

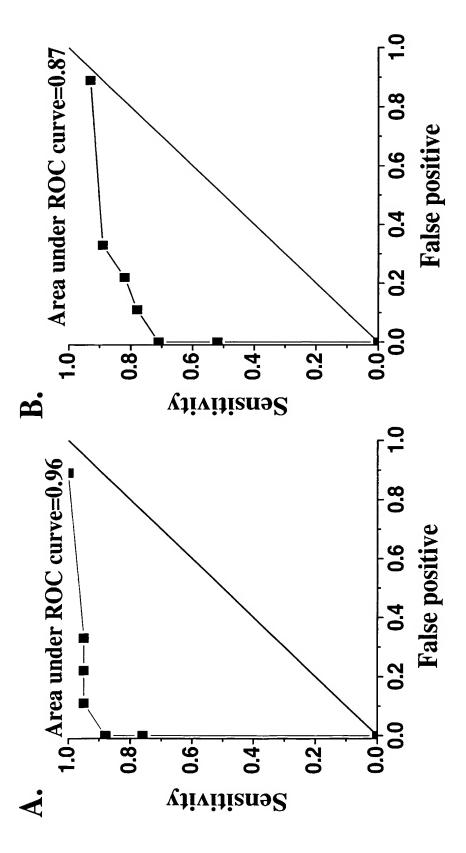


Figure 12

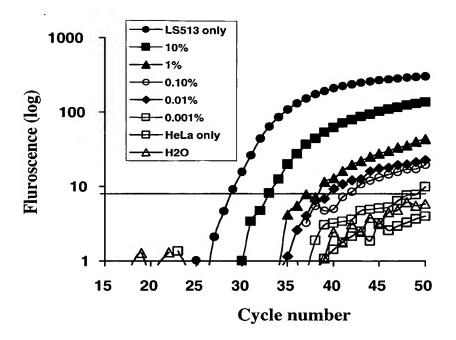


Figure 13

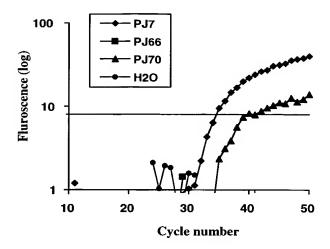


Figure 14

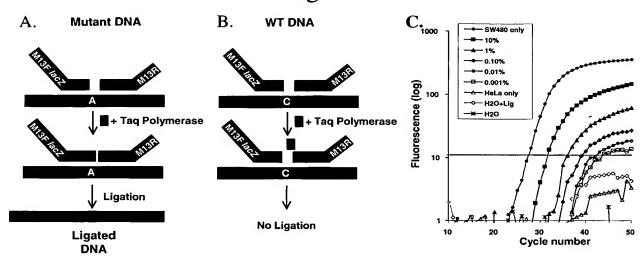


Figure 15

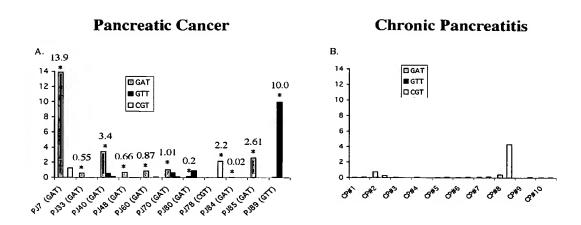


Figure 16

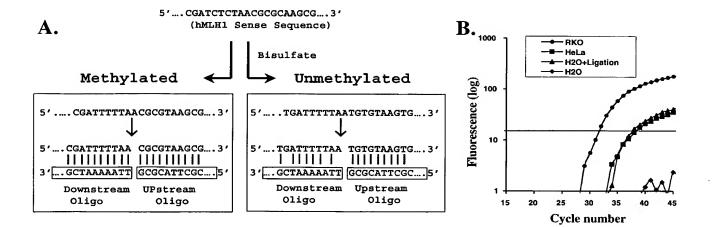


Figure 17

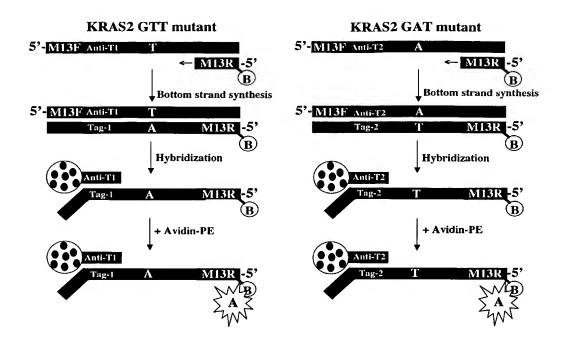


Figure 18

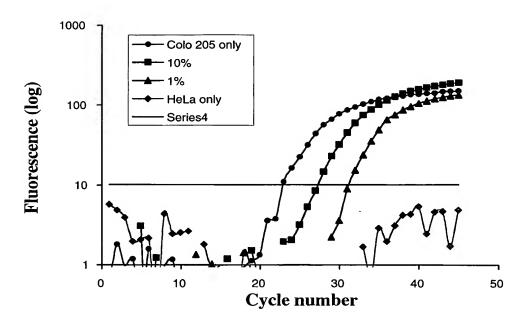
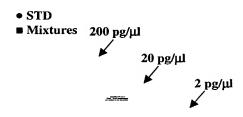


Figure 19

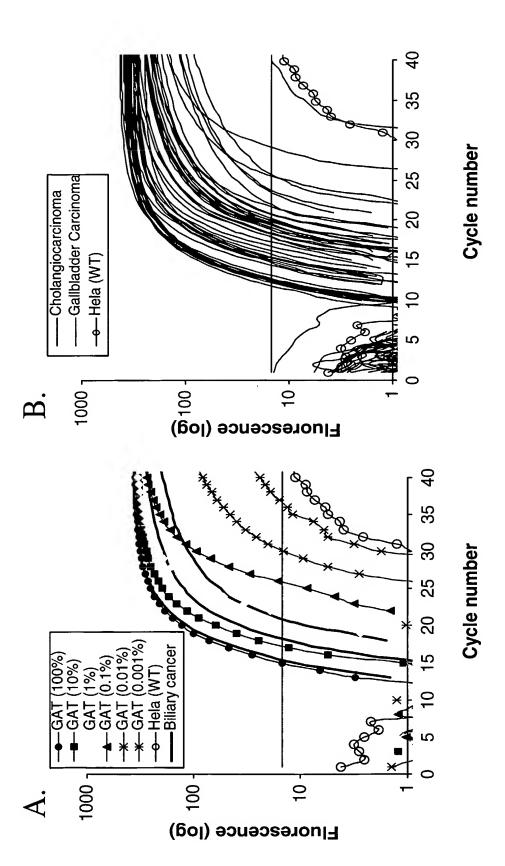
A. Standard curve for quantification



B. % heteroplasmy in the mixtures

Sample #	Heteroplasmy (%)
A	55.3
В	18.7
С	3.3
D	55.7
E	1.1
F	27.5
G	4.02
Н	25.6

Figure 20



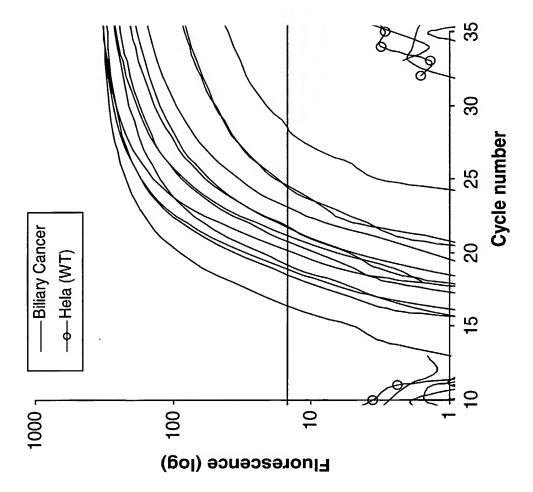
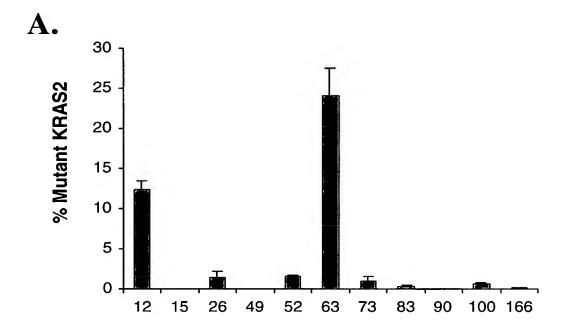


Figure 21

WO 2005/092038 PCT/US2005/009391 25/31

Figure 22



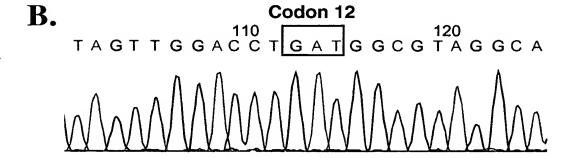
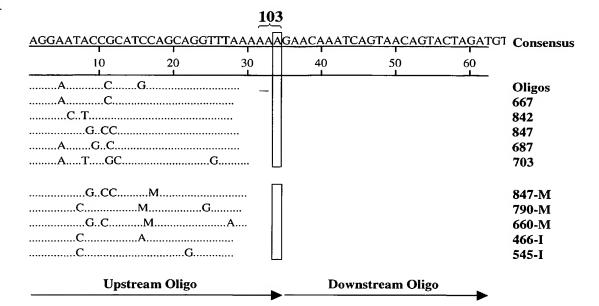
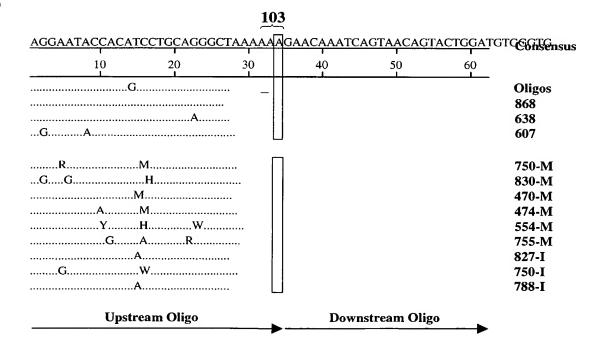


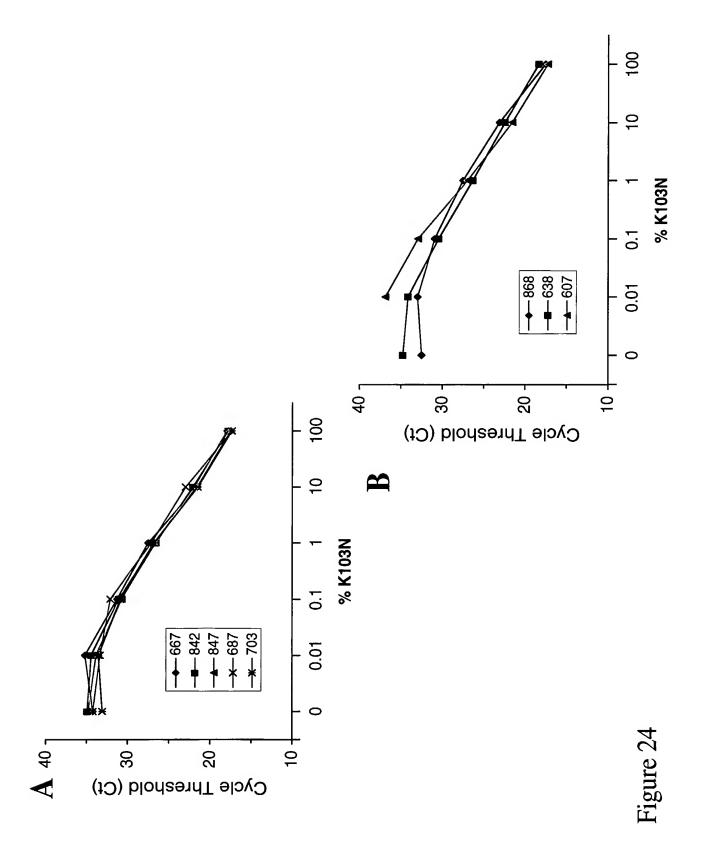
Figure 23

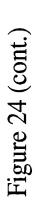
A

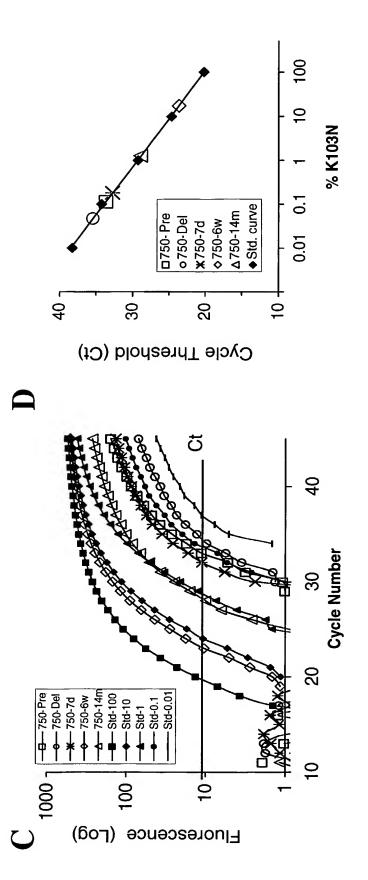


B









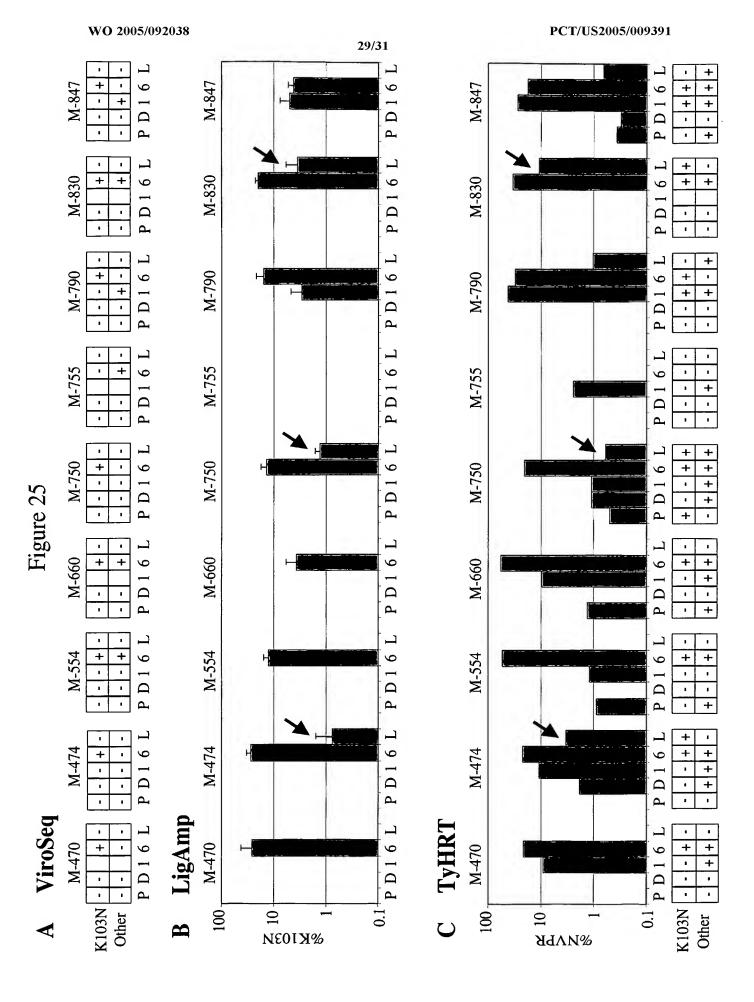


Figure 26

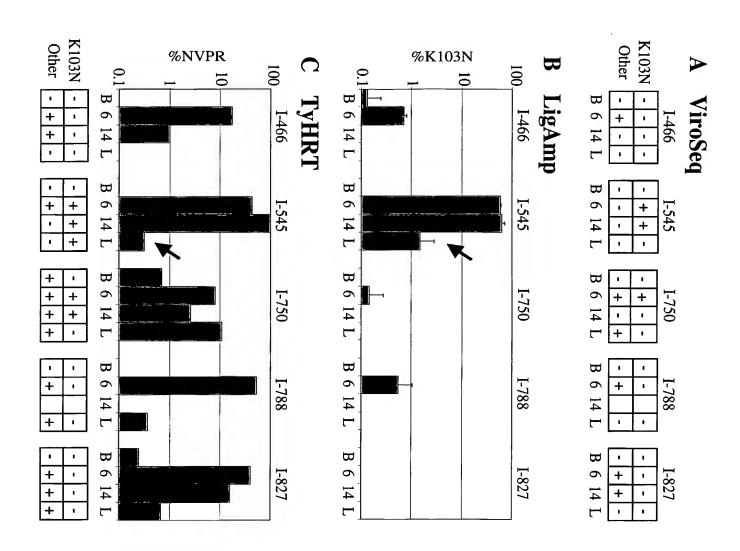
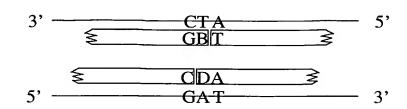
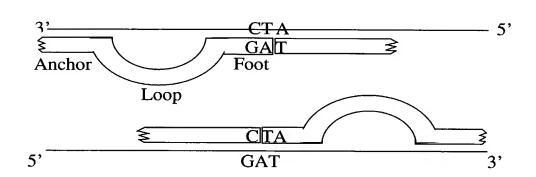


Figure 27

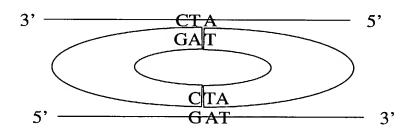
A.



B.



C.



D.

